Demultiplexing Report

Angela Crabtree

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Distribution of Read Pairs in Demultiplexed Bins

Inputs for the python script, demux.py, were 4 FASTQ files (R1, R2, R3, R4), one tab-delimited text file with barcode id and sequence information, and a user-specified q-score cutoff. The cutoff determined which reads failed to have good-quality barcode sequences and were thus relegated to the "junk" bin. Note that this cutoff does NOT apply to the sequencing coverage of the reads themselves.

Here are the command-line arguments used:

```
## ./demux_err.py
## -1 /projects/bgmp/shared/2017_sequencing/1294_S1_L008_R1_001.fastq.gz
## -2 /projects/bgmp/shared/2017_sequencing/1294_S1_L008_R2_001.fastq.gz
## -3 /projects/bgmp/shared/2017_sequencing/1294_S1_L008_R3_001.fastq.gz
## -4 /projects/bgmp/shared/2017_sequencing/1294_S1_L008_R4_001.fastq.gz
## -b /projects/bgmp/shared/2017_sequencing/indexes.txt
## -q 20
## -0 REAL-output
```

If barcodes in FASTQ files did not match barcodes in the barcode info file, they were also binned in "junk". If both forward and reverse barcodes for a read pair matched those in info file, but were not the same, read swapping occurred and read pairs were binned in "swap" bins. If barcodes matched, they were binned in the appropriate index bin. Counts were recorded and written to a csv output file.

Here are the counts of read pairs binned:

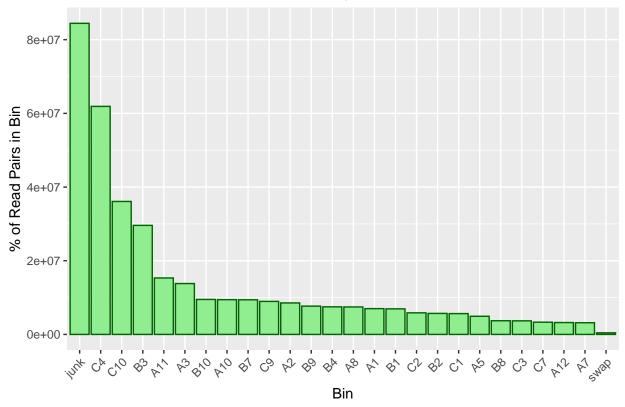
```
##
       bin rpcount
## 1
      junk 84433220 23.2440410
## 2
      swap
             432046
                      0.1189401
## 3
        В1
            6941570
                     1.9109793
## 4
        A5
            4926263
                      1.3561754
## 5
        C1
            5653676
                      1.5564286
## 6
        B9
            7674791
                      2.1128314
## 7
        C9
            8964532
                     2.4678906
## 8
        СЗ
            3688632
                      1.0154618
## 9
        B3 29599418
                     8.1485710
## 10
        C4 61891888 17.0385256
       A11 15322832
                     4.2182986
## 11
## 12
        C7
            3318839
                      0.9136597
## 13
        B2
            5702479
                      1.5698638
##
  14
        Α1
            7001570
                     1.9274970
## 15
            9400229
                     2.5878358
        A3 13809544 3.8016980
## 16
```

```
## 17
       В4
           7492308 2.0625947
## 18
       A12
           3225068 0.8878450
       C10 36089262
                     9.9351924
##
  20
        A2
            8546398
                     2.3527804
##
  21
       C2
            5858542
                     1.6128272
  22
            9433812 2.5970810
      A10
  23
       В8
            3715385
                     1.0228268
  24
                     0.8739839
##
        A7
            3174718
##
  25
       B10
            9490524
                     2.6126935
## 26
           7459189 2.0534772
        8A
```

bin = index id or bin name; rpcount = read pair count; rpp = read pair percent of total

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Other Stats

Percent of Junk Reads:

23.24404 %

Percent of Swapped Reads:

0.1189401 %

Summary statistics:

##	bin	rpcount	rpp
##	A1 : 1	Min. : 432046	Min. : 0.1189
##	A10 : 1	1st Qu.: 5108116	1st Qu.: 1.4062
##	A11 : 1	Median : 7475748	Median : 2.0580
##	A12 : 1	Mean :13971028	Mean : 3.8462
##	A2 : 1	3rd Qu.: 9476346	3rd Qu.: 2.6088
##	A3 : 1	Max. :84433220	Max. :23.2440
##	(Other):20		